

Borrelia recurrentis の性状と回帰熱
ボレリア、ライム病関連ボレリア、
その他ボレリア種との比較

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Borrelia recurrentis Characterization and Comparison with
Relapsing-Fever, Lyme-Associated, and Other *Borrelia* spp.

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ABSTRACT *Borrelia recurrentis*, the cause of louse-borne relapsing fever, has until recently been considered noncultivable, which has prevented characterization of this spirochete. We successfully cultivated 18 strains from patients with louse-borne relapsing fever and present the initial characterization of these isolates. Electron microscopy revealed spirochetal cells with pointed ends, an average wavelength of $1.8\ \mu\text{m}$, an amplitude of $0.8\ \mu\text{m}$, and 8 to 10 periplasmic flagella. The G+C ratio was 28.4 mol%. Whole DNA-DNA hybridizations showed similarity between the isolates of *B. recurrentis* but not with *Borrelia hermsii*, *Borrelia parkeri*, *Borrelia turicatae*, or the Lyme-associated borreliae. Sequencing studies of both the flagellin and 16S rRNA genes revealed that the greatest similarity was between *B. recurrentis* and *Borrelia duttonii*. Analysis of the sodium dodecyl sulfate-polyacrylamide gel electrophoresis profiles of strains revealed four groups based on the position of a major protein band (one of the groups showed some heterogeneity and was subdivided into four subgroups). Pulsed-field gel electrophoresis revealed five distinct patterns.

抄録 シラミ媒介性回帰熱の起因菌である *Borrelia recurrentis* は、これまで培養が不

可能であったことからその性状はあまり知られていない。我々はシラミ媒介性回帰熱の患者から18株のボレリアの培養に成功し、これら分離株の性状について検討した。電子顕微鏡像、G+C含量、DNA 相同性、鞭毛遺伝子あるいは16S rRNA 遺伝子の塩基配列、SDS-PAGE、PFGEによる解析を行い、回帰熱ボレリア、ライム病関連ボレリア、その他ボレリア属に属する種との比較を行った。

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